SEQUENCE LISTING

<110> HEIN, MICH B. HIATT, ANDREW C. FITCHEN, JOHN H. <120> NOVEL EPITHELIAL TISSUE TARGETING AGENT <130> 040989/283662 <140> 09/005,318 <141> 1998-01-09 <150> 08/782,481 <151> 1997-01-10 <150> 09/005,167 <151> 1998-01-09 <160> 140 <170> PatentIn Ver. 2.1 <210> 1 <211> 137 <212> PRT <213> Homo sapiens <400> 1 Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg Glu 35 Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Pro Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp 75 Asn Gln Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser 85

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Leu Thr Pro Asp Ala Cys Tyr Pro Asp 130 135

115

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Arg Ile Thr Ser Arg Ile Ile Pro Ser Ala Glu Asp Pro Ser Gln Asp
20 25 30

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35 40 45

Asn Ile Ser Asp Pro Thr Ser Pro Met Arg Thr Lys Pro Val Tyr His 50 55 60

Leu Ser Asp Leu Cys Lys Lys Cys Asp Thr Thr Glu Val Glu Leu Glu 65 70 75 80

Asp Gln Val Val Thr Ala Ser Gln Ser Asn Ile Cys Asp Ser Asp Ala 85 90 95

Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Asn Arg Val 100 105 110

Lys Leu Ser Tyr Arg Gly Gln Thr Lys Met Val Glu Thr Ala Leu Thr
115 120 125

Pro Asp Ser Cys Tyr Pro Asp 130 135

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<211> 137

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<213> Oryctolagus cuniculus

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Val Glu Arg Asn Ile Arg Ile Val Val Pro Leu Asn Asn Arg Glu Asn 35 40 45

Ile Ser Asp Pro Thr Ser Pro Leu Arg Arg Asn Pro Val Tyr His Leu 50 55 60

Ser Asp Val Cys Lys Lys Cys Asp Pro Val Glu Val Glu Leu Glu Asp 65 70 75 80

Gln Val Val Thr Ala Thr Gln Ser Asn Ile Cys Asn Glu Asp Asp Gly
85 90 95

Val Pro Glu Thr Cys Tyr Met Tyr Asp Arg Asn Lys Cys Tyr Thr Thr 100 105 110

Met Val Pro Leu Arg Tyr His Gly Glu Thr Lys Met Val Gln Ala Ala 115 120 125

Leu Thr Pro Asp Ser Cys Tyr Pro Asp 130 135

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20 25 30

Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Thr Arg Glu Asn 35 40 45

Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Glu Pro Lys Tyr Asn Leu 50 55 60

Ala Asn Leu Cys Lys Lys Cys Asp Pro Thr Glu Ile Glu Leu Asp Asn 65 70 75 80

Gln Val Phe Thr Ala Ser Gln Ser Asn Ile Cys Pro Asp Asp Tyr 85 90 95

Ser Glu Thr Cys Tyr Thr Tyr Asp Arg Asn Lys Cys Tyr Thr Thr Leu 100 105 110

Val Pro Ile Thr His Arg Gly Val Thr Arg Met Val Lys Ala Thr Leu 115 120 125

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Glu Arg Asn Ile Gln Ile Thr Ile Pro Thr Ser Ser Arg Met Xaa Ile 35 40 45

Ser Asp Pro Tyr Ser Pro Leu Arg Thr Gln Pro Val Tyr Asn Leu Trp 50 55 60

Asp Ile Cys Gln Lys Cys Asp Pro Val Gln Leu Glu Ile Gly Gly Ile
65 70 75 80

Pro Val Leu Ala Ser Gln Pro Xaa Xaa Ser Xaa Pro Asp Asp Glu Cys 85 90 95

Tyr Thr Thr Glu Val Asn Phe Lys Lys Lys Val Pro Leu Thr Pro Asp 100 105 110

Ser Cys Tyr Glu Tyr Ser Glu 115

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Pro Leu Lys Asn Arg Gly Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg 35 40 45

Asn Gln Pro Val Tyr His Leu Ser Pro Ser Cys Lys Lys Cys Asp Pro 50 60

Tyr Glu Asp Gly Val Val Thr Ala Thr Glu Thr Asn Ile Cys Tyr Pro 65 70 75 80

Asp Gln Gly Val Pro Gln Ser Cys Arg Asp Tyr Cys Pro Glu Leu Asp

85 90 95

Arg Asn Lys Cys Tyr Thr Val Leu Val Pro Pro Gly Tyr Thr Gly Glu
100 105 110

Thr Lys Met Val Gln Asn Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp 115 120 125

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gct cgt att act tct aga atc atc cgt agc tca gag gac cca aat gaa 96
Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser Glu Asp Pro Asn Glu
15 20 25 30

gat ata gtc gaa cgt aac atc cgt atc atc gtc cca ctg aat aac cgg 144
Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg
35 40 45

gag aat atc tca gat cct aca agt ccg ttg cgc aca cgc ttc gta tac 192
Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr
50 55 60

cac ctg tca gat ctg tgt aag aag tgt gat cca aca gag gta gag ctg 240 His Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu 65 70 75

gac aat cag ata gtc act gcg act caa agc aac att tgc gat gag gac 288
Asp Asn Gln Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp
80 85 90

agc gct aca gaa acc tgc agc acc tac gat agg aac aaa tgc tac acg 336 Ser Ala Thr Glu Thr Cys Ser Thr Tyr Asp Arg Asn Lys Cys Tyr Thr 95 100 105 110

gcc gtg gtt ccg ctc gtg tat ggt gga gag aca aaa atg gtg gaa act 384 Ala Val Val Pro Leu Val Tyr Gly Gly Glu Thr Lys Met Val Glu Thr 115 120 125

gcc ctt acg ccc gat gca tgc tat ccg gac tgaattc Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp 130 135	421		
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tca gag gac cca aat gaa gat ata gtc gaa cgt aac atc cgt atc atc Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile 20 25 30	96		
gtc cca ctg aat aac cgg gag aat atc tca gat cct aca agt ccg ttg Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu 35 40 45	144		
cgc aca cgc ttc gta tac cac ctg tca gat ctg tgt aag aag gat gag Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Asp Glu 50 55 60	192		
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caaaaatggt ggaaactgcc cttacgcccg atgcatgcta ccctgactg
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Asp Asn Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser
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tca gag gac cca aat gaa gat ata gtc gaa cgt aac atc cgt atc atc
Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile
             20
                                  25
gtc cca ctg aat aac cgg gag aat atc tca gat cct aca agt ccg ttg
                                                                   144
Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu
         35
                             40
cgc aca cgc ttc gta tac cac ctg tca gat ctg tgt aag aag tgt gat
                                                                   192
Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp
     50
                         55
cca aca gag gta gag ctg gac aat cag ata gtc act gcg act caa agc
                                                                   240
Pro Thr Glu Val Glu Leu Asp Asn Gln Ile Val Thr Ala Thr Gln Ser
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70 75 80 65 aac att tgc gat gag gac agc gct aca gaa acc tgc tac tga 282 Asn Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr $\,\,$ * 85 90 286 attc <210> 14 <211> 105 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(105) <400> 14 48 gat ctg tgt aag aag tgt gat cca aca gag gta gag ctg gac aat cag Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu Asp Asn Gln ata gtc act gcg act caa agc aac att tgc gat gag gac agc gct aca 96 Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp Ser Ala Thr 25 gaa acc tgc 105 Glu Thr Cys 35 <210> 15 <211> 61 <212> DNA <213> Homo sapiens <400> 15 gatcaggaag atgaacgtat tgttctggtt gacaacaagt gcaagtgtgc tcgtattact 60 61 <210> 16 <211> 198 <212> DNA <213> Homo sapiens <400> 16 gcgatgacga cgataaggcc caaacggaga cctgtactgt tgcgcctcgt gaacggcaaa 60 actgoggatt cccgggagta acaccctctc agtgcgctaa taaaggctgc tgttttgatg 120 acacggtacg gggcgttccg tggtgcttct accccaatac aattgacgtt ccgcctgaag 180 aagagtgcga gttttaag 198

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<213> Homo sapiens

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15 20 25 30

Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg
35 40 45

Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr
50 55 60

His Leu Ser Asp Leu Cys Lys Lys Cys Asp Pro Thr Glu Val Glu Leu 65 70 75

Asp Asn Gln Ile Val Thr Ala Thr Gln Ser Asn Ile Cys Asp Glu Asp 80 85 90

Ser Ala Thr Glu Thr Cys Ser Thr Tyr Asp Arg Asn Lys Cys Tyr Thr 95 100 105 110

Ala Val Val Pro Leu Val Tyr Gly Gly Glu Thr Lys Met Val Glu Thr 115 120 125

Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp 130 135

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<212> PRT

<213> Homo sapiens

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Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile 20 25 30

Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu 35 40 45

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Asp Ser Ala Thr Glu Thr Cys 65 70

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<213> Homo sapiens
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Arg Asn Ile Arg Ile Ile Val Pro Leu Asn Asn Arg Glu Asn Ile Ser
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Asp Pro Thr Ser Pro Leu Arg Thr Arg Phe Val Tyr His Leu Ser Asp
Leu
<210> 20
<211> 12
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<213> Homo sapiens
<400> 20
Asp Gln Lys Cys Lys Cys Ala Arg Ile Thr Ser Arg
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Asp Leu Cys Lys Lys Asp Glu Asp Ser Ala Thr Glu Thr Cys
<210> 22
<211> 36
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<213> Homo sapiens
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Tyr Gly Gly Glu Thr Lys Met Val Glu Thr Ala Leu Thr Pro Asp Ala
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Cys Tyr Pro Asp
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<213> Homo sapiens

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Ser Glu Asp Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile
20 25 30

Val Pro Leu Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu 35 40 45

Arg Thr Arg Phe Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp 50 55 60

Pro Thr Glu Val Glu Leu Asp Asn Gln Ile Val Thr Ala Thr Gln Ser 65 70 75 80

Asn Ile Cys Asp Glu Asp Ser Ala Thr Glu Thr Cys Tyr
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20 25 30

Glu Thr Cys

<210> 25

<211> 22

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Ala Arg Ile Thr Ser Arg
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Pro Arg Glu Arg Gln Asn Cys Gly Phe Pro Gly Val Thr Pro Ser Gln
Cys Ala Asn Lys Gly Cys Cys Phe Asp Asp Thr Val Arg Gly Val Pro
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Trp Cys Phe Tyr Pro Asn Thr Ile Asp Val Pro Pro Glu Glu Glu Cys
     50
                         55
Glu Phe
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tttctgtagc gctgtcctca tcgcaaatgt tgctttgagt cgcagtgact atctgattgt 180
ccagetetae etetgttgga teacaettet tacacagate tgacaggtgg tatacgaage 240
gtgtgcgcaa cggacttgta ggatctgaga tattctcccg gttattcagt gggacgatga 300
tacggatgtt acgttcgact atatcttcat ttgggtcctc tgagctacgg atgattctag 360
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gacgatgata cggatgttac gttcgactat atcttcattt gggtcctctg agctacggat 180
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ttactcccgg gaatccgcag ttttgccgtt cacgaggcgc aacagtacag gtctccgttt 180
gggccttatc gtcgtcatcg ctgca
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<210> 39
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tac atc tat gcg gat ccg agc tcg agt gct ctagatctgc agctggtacc
                                                                    98
Tyr Ile Tyr Ala Asp Pro Ser Ser Ser Ala
             20
                                  25
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atggaattcg aagcttggag tcgactctgc tga
<210> 43
<211> 26
<212> PRT
<213> Homo sapiens
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<211> 61
<212> DNA
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<211> 61
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       oligonucleotide
 <400> 58
 acggacttgt aggatctgag atattctccc ggttattcag tgggacgat
                                                                     49
 <210> 59
 <211> 49
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       oligonucleotide
 <400> 59
 acggacttgt aggatctgag atgtgctccc ggttattcag tgggacgat
                                                                    49
 <210> 60
 <211> 44
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: Synthetic
       oligonucleotide
 <400> 60
 atcctacaag tccgttgcgc acacgcttcg tataccacct gtca
                                                                    44
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<210> 61
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 61
gatctgacag gtggtatacg aagcgtgtgc gca
                                                                    33
<210> 62
<211> 60
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 62
gatctgtgta agaagtgtga tccaacagag gtagagctgg acaatcagat agtcactgca 60
<210> 63
<211> 44
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 63
gatctgtgta agaaggatga ggacagcgct acagaaacct gctg
                                                                    44
<210> 64
<211> 44
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 64
aattcagcag gtttctgtag cgctgtcctc atccttctta caca
                                                                    44
<210> 65
<211> 62
<212> DNA
<213> Artificial Sequence
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<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 65
gatctgtgta agaaggatga ggacagcgct acagaaacct gctacgagaa ggatgagctg 60
<210> 66
<211> 62
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 66
aattcacage teateetteg egtegeaggt ttetgtageg etgteeteat cettettaca 60
                                                                   62
<210> 67
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 67
qatctqtqta aqaaqtctqa tatcgatqaa gattccgcta cagaaacctg cagcacatg 59
<210> 68
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 68
aattcatgtg ctgcaggttt ctgtagcgga atcttcatcg atatcagact tcttacaca 59
<210> 69
<211> 64
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
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oligonucleotide

<400> gatctg ctaa	69 gtcta agaagtctga tatcgatgaa gattacagat tcttcagact atagctactt	60 64
<210><211><212><212><213>	30	
<220> <223>	Description of Artificial Sequence: Synthetic oligonucleotide	
<400> aatctt	70 tcatc gatatcagac ttcttagaca	30
<210> <211> <212> <213>	64	
<220> <223>	Description of Artificial Sequence: Synthetic oligonucleotide	
<400> gatcto	71 ggtta agaagtctga tatcgatgaa gattaccaat tcttcagact atagctactt	60 64
<210><211><211><212><213>	30	
<220> <223>	Description of Artificial Sequence: Synthetic oligonucleotide	
<400> aatctt	72 tcatc gatatcagac ttcttaacca	30
<210><211><211><212><213>	41	
<220> <223>	Description of Artificial Sequence: Synthetic oligonucleotide	

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<400> 73
                                                                    41
attgtccagc tctacctctg ttggatcaca cttcttacac a
<210> 74
<211> 46
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 74
actcaaagca acatttgcga tgaggacagc gctacagaaa cctgca
                                                                   46
<210> 75
<211> 57
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 75
ggtttctgta gcgctctgct catcgcaaat gttgctttga gtcgcagtga ctatctg
<210> 76
<211> 59
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 76
gcacctacga taggaacaaa tgctacacgg ccgtggttcc gctcgtgtat ggtggagag 59
<210> 77
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
gagcggaacc acggccgtgt agcatttgtt cctatcgtag gtgctgca
                                                                   48
<210> 78
```

```
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 78
acaaaaatgg tggaaactgc ccttacgccc gatgcatgct atccggactg
                                                                    50
<210> 79
<211> 69
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
aattcagtcc ggatagcatg catcgggcgt aagggcagtt tccaccattt ttgtctctcc 60
                                                                    69
accatacac
<210> 80
<211> 62
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 80
acaaaaatgg tggaaactgc ccttacgccc gatgcatgct atccggacaa ggatgaattg 60
<210> 81
<211> 81
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 81
aattcacaat tcatccttgt ccggatagca tgcatcgggc gtaagggcag tttccaccat 60
ttttgtctct ccaccataca c
<210> 82
<211> 88
<212> DNA
```

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 82
gatcaggtcg ctgccatcca agacccgagg ctgttcgccg aagagaaggc cgtcgctgac 60
tccaagtgca agtgtgctcg tattactt
<210> 83
<211> 88
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 83
ctagaagtaa tacgagcaca cttgcacttg gagtcagcga cggccttctc ttcggcgaac 60
agcctcgggt cttggatggc agcgacct
<210> 84
<211> 34
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 84
                                                                    34
tggtacgaat tccaggtsma rctgcagsag tcrg
<210> 85
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 85
                                                                    27
acagatatcg ggatttctcg cagactc
<210> 86
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
```

```
<400> 86
                                                                    28
acagaatatc gtcaacacct tcccaccc
<210> 87
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 87
                                                                    30
acaaagcttt tatttacccg acagacggtc
<210> 88
<211> 35
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 88
gtccccctc gagcgayaty swgmtsaccc artct
                                                                    35
<210> 89
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 89
acactgcagc agttggtgca gcatcagc
                                                                    28
<210> 90
<211> 53
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 90
ctgcaggaag cggaagcgga ggaagcggaa gcggaggaag cggaagcgaa ttc
                                                                    53
<210> 91
<211> 47
<212> DNA
```

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Linker
      complement
<400> 91
                                                                    47
cettegeett egeeteette geettegeet eettegeett egettaa
<210> 92
<211> 76
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Signal peptide
<400> 92
acaggateca tggaaacece agegeagett etetteetee tgetaetetg geteecaaga 60
taccaccgga cccggg
<210> 93
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 93
                                                                    33
tggtacagat ctaggtsmar ctgcagsagt crg
<210> 94
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 94
                                                                    28
acaggaattc aattttcttg tccacctt
<210> 95
<211> 29
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 95
gttctagaga yatyswgmts acccartct
                                                                    29
```

```
<210> 96
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 96
                                                                    28
acaccgcggc agttggtgca gcatcagc
<210> 97
<211> 75
<212> DNA
<213> Homo sapiens
<400> 97
acaggateca tggaaacccc agegeagett etetteetee tgetaetetg geteecagat 60
accaccggaa gatct
<210> 98
<211> 75
<212> DNA
<213> Homo sapiens
<400> 98
acaactagta tggaaacccc agcgcagctt ctcttcctcc tgctactctg gctcccagat 60
accaccggat ctaga
                                                                    75
<210> 99
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Linker peptide
Val Ala Val Gln Ser Ala Gly Thr Pro Ala Ser Gly Ser
                  5
<210> 100
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Nuclear
      targeting sequence
```

```
<400> 100
Cys Ala Ala Pro Lys Lys Lys Arg Lys Val
                  5
<210> 101
<211> 22
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Nuclear
      targeting sequence
<400> 101
Cys Ala Ala Lys Arg Pro Pro Ala Ala Ile Lys Lys Ala Ala Ala Gly
  1
                  5
Gln Ala Lys Lys Lys
             20
<210> 102
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Intracellular
      targeting signal
<400> 102
His Asp Glu Leu
<210> 103
<211> 77
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 103
gcgatgacga cgataaggcc caaacggaga cctgtactgt tgcgcctcgt gaacggcaaa 60
actgcggatt cccggga
<210> 104
<211> 66
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
```

oligonucleotide

```
<400> 104
gttttgccgt tcacgaggcg caacagtaca ggtctccgtt tgggccttat cgtcgtcatc 60
gctgca
<210> 105
<211> 72
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 105
qtaacaccct ctcaqtqcqc taataaaggc tgctgttttg atgacacggt acggggcgtt 60
                                                                    72
ccgtggtgct tc
<210> 106
<211> 72
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 106
gccccgtacc gtgtcatcaa aacagcagcc tttattagcg cactgagagg gtgttactcc 60
cgggaatccg ca
                                                                    72
<210> 107
<211> 49
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
<400> 107
taccccaata caattgacgt tccgcctgaa gaagagtgcg agttttaag
                                                                    49
<210> 108
<211> 68
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      oligonucleotide
```

```
aattettaaa aetegeacte ttetteagge ggeaagteaa ttgtattggg gtagaageac 60
cacggaac
<210> 109
<211> 7
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Linker peptide
<400> 109
Pro Leu Gly Ile Ile Gly Gly
<210> 110
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Linker peptide
<400> 110
Ile Ile Gly Gly
  1
<210> 111
<211> 30
<212> PRT
<213> Homo sapiens
<400> 111
Val Arg Asp Gln Ala Gln Glu Asn Arg Ala Ser Gly Asp Ala Gly Ser
Ala Asp Gly Gln Ser Arg Ser Ser Ser Lys Val Leu Phe
                                 25
<210> 112
<211> 25
<212> PRT
<213> Homo sapiens
<400> 112
Val Pro Ser Thr Pro Pro Thr Pro Ser Pro Ser Thr Pro Pro Thr Pro
                                     10
Ser Pro Ser Cys Cys His Pro Arg Leu
```

20

<400> 108

```
<210> 113
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Illustrative
      peptide
<400> 113
Glu Gln Lys Leu Ile Ser Glu Asp Leu
<210> 114
<211> 68
<212> PRT
<213> Homo sapiens
<400> 114
Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Arg Ser Ser Glu Asp
                                     10
Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu
            20
Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Arg
Pro Val Tyr His Leu Ser Asp Leu Cys Lys Lys Asp Glu Asp Ser Ala
                        55
                                             60
Thr Glu Thr Cys
65
<210> 115
<211> 12
<212> PRT
<213> Homo sapiens
<400> 115
Cys Lys Cys Ala Arg Asp Ser Asp Ala Glu Thr Cys
                 5
<210> 116
<211> 69
<212> PRT
<213> Homo sapiens
<400> 116
Cys Met Cys Thr Arg Val Thr Ser Arg Ile Ile Pro Ser Thr Glu Asp
                                    10
Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Val Val Pro Leu
Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Arg Asn
                            40
Pro Val Tyr His Leu Ser Asp Val Cys Lys Lys Asn Glu Asp Asp Gly
```

```
60
    50
                        55
Val Pro Glu Thr Cys
65
<210> 117
<211> 68
<212> PRT
<213> Homo sapiens
<400> 117
Cys Gln Cys Val Arg Ile Thr Ser Arg Ile Ile Arg Asp Pro Asp Asn
 1
                                    10
Pro Ser Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu
            20
Asn Thr Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Glu
                            40
Pro Lys Tyr Asn Leu Ala Asn Leu Cys Lys Lys Pro Asp Asp Tyr
                        55
Ser Glu Thr Cys
<210> 118
<211> 67
<212> PRT
<213> Homo sapiens
<220>
<221> VARIANT
<222> 37, 60, 62
<223> Xaa = Any Amino Acid
<400> 118
Cys Lys Cys Val Lys Ile Ser Ser Arg Phe Val Pro Ser Thr Glu Arg
                                    10
Pro Gly Glu Glu Ile Leu Glu Arg Asn Ile Gln Ile Thr Ile Pro Thr
                                25
Ser Ser Arg Met Xaa Ile Ser Asp Pro Tyr Ser Pro Leu Arg Thr Gln
Pro Val Tyr Asn Leu Trp Asp Ile Cys Gln Lys Xaa Ser Xaa Pro Asp
    50
                        55
Asp Glu Cys
65
<210> 119
<211> 69
<212> PRT
<213> Homo sapiens
<400> 119
Cys Met Cys Thr Arg Val Thr Ala Arg Ile Arg Gly Thr Arg Glu Asp
1
Pro Asn Glu Asp Ile Val Glu Arg Tyr Ile Arg Ile Asn Val Pro Leu
                                25
Lys Asn Arg Gly Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Asn Gln
```

```
40
Pro Val Tyr His Leu Ser Pro Ser Cys Lys Lys Tyr Pro Asp Gln Gly
                        55
Val Pro Gln Ser Cys
<210> 120
<211> 87
<212> PRT
<213> Homo sapiens
<400> 120
Cys Lys Cys Ala Arg Ile Thr Ser Arg Ile Ile Pro Ser Ala Glu Asp
                                    10
Pro Ser Gln Asp Ile Val Glu Arg Asn Val Arg Ile Ile Val Pro Leu
                                25
Asn Ser Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Met Arg Thr Lys
                                                 45
Pro Val Tyr His Leu Ser Asp Leu Cys Lys Lys Cys Asp Thr Thr Glu
Val Glu Leu Glu Asp Gln Val Val Thr Ala Ser Gln Ser Asn Ile Cys
                    70
                                        75
Asp Ser Asp Ala Glu Thr Cys
                85
<210> 121
<211> 90
<212> PRT
<213> Homo sapiens
<400> 121
Cys Met Cys Thr Arg Val Thr Ser Arg Ile Ile Pro Ser Thr Glu Asp
                                    10
Pro Asn Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Val Val Pro Leu
Asn Asn Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Arg Asn
                            40
Pro Val Tyr His Leu Ser Asp Val Cys Lys Cys Asp Pro Val Glu
                      . 55
                                            60
Val Glu Leu Glu Asp Gln Val Val Thr Ala Thr Gln Ser Asn Ile Cys
                    70
Asn Glu Asp Asp Gly Val Pro Glu Thr Cys
                85
<210> 122
<211> 89
<212> PRT
<213> Homo sapiens
<400> 122
Cys Gln Cys Val Arg Ile Thr Ser Arg Ile Ile Arg Asp Pro Asp Asn
                                    10
Pro Ser Glu Asp Ile Val Glu Arg Asn Ile Arg Ile Ile Val Pro Leu
```

25

20

Asn Thr Arg Glu Asn Ile Ser Asp Pro Thr Ser Pro Leu Arg Thr Glu 35

Pro Lys Tyr Asn Leu Ala Asn Leu Cys Lys Lys Cys Asp Pro Thr Glu 50

Ile Glu Leu Asp Asn Gln Val Phe Thr Ala Ser Gln Ser Asn Ile Cys 65

Pro Asp Asp Asp Tyr Ser Glu Thr Cys 85

<210> 123 <211> 85 <212> PRT <213> Homo sapiens

<220>

<221> VARIANT <222> 37, 82, 83, 85

<223> Xaa = Any Amino Acid

<400> 123

 Cys
 Lys
 Cys
 Val
 Lys
 Ile
 Ser
 Ser
 Arg
 Phe
 Val
 Pro
 Ser
 Thr
 Glu
 Arg

 Pro
 Gly
 Glu
 Glu
 Ile
 Leu
 Glu
 Arg
 Asn
 Ile
 Gln
 Ile
 Pro
 Thr
 Ile
 Pro
 Thr
 Ile
 Pro
 Thr
 Gln
 Ile
 Ile

85

<210> 124 <211> 84 <212> PRT <213> Homo sapiens

<400> 124

 Cys
 Thr
 Arg
 Val
 Thr
 Ala
 Arg
 Ile
 Arg
 Gly
 Thr
 Arg
 Gly
 Thr
 Arg
 Gly
 Thr
 Arg
 Gly
 Thr
 Arg
 Ile
 Arg
 Arg
 Pro
 Leu
 Arg
 Arg
 Gln
 Pro
 Val

 Arg
 A

<210> 125 <211> 12

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<212> PRT
<213> Homo sapiens
<400> 125
Gln Glu Asp Glu Arg Ile Val Leu Val Asp Asn Lys
               5
<210> 126
<211> 12
<212> PRT
<213> Homo sapiens
<400> 126
Gln Asp Glu Asn Glu Arg Ile Val Val Asp Asn Lys
                5
<210> 127
<211> 11
<212> PRT
<213> Homo sapiens
<400> 127
Asp Asp Glu Ala Thr Ile Leu Ala Asp Asn Lys
               5
<210> 128
<211> 11
<212> PRT
<213> Homo sapiens
<400> 128
Asp Asp Glu Ala Thr Ile Leu Ala Asp Asn Lys
               5
<210> 129
<211> 10
<212> PRT
<213> Homo sapiens
Glu Gln Glu Tyr Ile Leu Ala Asn Asn Lys
                5
<210> 130
<211> 7
<212> PRT
<213> Homo sapiens
<400> 130
Tyr Thr Tyr Asp Arg Asn Lys
         5
```

```
<210> 131
<211> 7
<212> PRT
<213> Homo sapiens
<400> 131
Tyr Thr Tyr Asp Arg Asn Lys
                5
<210> 132
<211> 7
<212> PRT
<213> Homo sapiens
<400> 132
Tyr Met Tyr Asp Arg Asn Lys
<210> 133
<211> 7
<212> PRT
<213> Homo sapiens
<400> 133
Tyr Thr Tyr Asp Arg Asn Lys
<210> 134
<211> 11
<212> PRT
<213> Homo sapiens
<400> 134
Arg Asp Tyr Cys Pro Glu Leu Asp Arg Asn Lys
                5
<210> 135
<211> 29
<212> PRT
<213> Homo sapiens
<400> 135
Cys Tyr Thr Ala Val Val Pro Leu Val Tyr Gly Glu Thr Lys Met
                 5
                                    10
Val Glu Thr Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp
            20
<210> 136
<211> 29
<212> PRT
<213> Homo sapiens
```

```
<400> 136
Cys Tyr Thr Asn Arg Val Lys Leu Ser Tyr Arg Gly Gln Thr Lys Met
                                     10
                5
Val Glu Thr Ala Leu Thr Pro Asp Ser Cys Tyr Pro Asp
<210> 137
<211> 29
<212> PRT
<213> Homo sapiens
<400> 137
Cys Tyr Thr Thr Met Val Pro Leu Arg Tyr His Gly Glu Thr Lys Met
1
Val Gln Ala Ala Leu Thr Pro Asp Ser Cys Tyr Pro Asp
                                25
            20
<210> 138
<211> 29
<212> PRT
<213> Homo sapiens
<400> 138
Cys Tyr Thr Thr Leu Val Pro Ile Thr His Arg Gly Val Thr Arg Met
Val Lys Ala Thr Leu Thr Pro Asp Ser Cys Tyr Pro Asp
                                25
<210> 139
<211> 24
<212> PRT
<213> Homo sapiens
<400> 139
Cys Tyr Thr Thr Glu Val Asn Phe Lys Lys Lys Val Pro Leu Thr Pro
1
                                    10
Asp Ser Cys Tyr Glu Tyr Ser Glu
            20
<210> 140
<211> 29
<212> PRT
<213> Homo sapiens
<400> 140
Cys Tyr Thr Val Leu Val Pro Pro Gly Tyr Thr Gly Glu Thr Lys Met
1
                                    10
Val Gln Asn Ala Leu Thr Pro Asp Ala Cys Tyr Pro Asp
```